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3. The method of claim 2, wherein the set of training peptides include peptides having a binding affinity for MHC class I molecules.
4. The method of claim 3, wherein the peptides included in the set of training peptides have a binding affinity for mouse MHC class I K^b.
5. The method of claim 2, wherein the set of test peptides include peptides having a binding affinity for MHC class I molecules.
6. The method of claim 5, wherein the peptides included in the set of test peptides have a binding affinity for mouse MHC class I K^b.
7. The method of claims 1 or 2, wherein the ANN comprises a multi-layer perceptron ANN trained by back-propagation of error.

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8. A system for identifying relative binding motifs for peptide-like molecules, comprising:
- (a) means for training an artificial neural network (ANN) with a set of training peptide-like molecules, each of known sequence and binding affinity;
 - (b) means for applying to the ANN at least one test peptide-like molecule, each of known sequence but unknown binding affinity;
 - (c) means for analyzing each applied test peptide-like molecule using the ANN to predict a relative binding affinity for each test peptide-like molecule.
9. A system for identifying relative peptide binding motifs, comprising:
- (a) means for training an artificial neural network (ANN) with a set of training peptides, each of known binding affinity, each peptide comprising a sequence of amino acids, each amino acid being binary coded as having or lacking specific features generally characteristic of amino acids;
 - (b) means for applying to the ANN at least one test peptide, each of unknown binding affinity, each peptide comprising a sequence of amino acids, each amino acid being binary coded as having or lacking specific features generally characteristic of amino acids;
 - (c) means for analyzing each applied test peptide using the ANN to predict a relative binding affinity for each test peptide.

10. The system of claim 9, wherein the set of training peptides include peptides having a binding affinity for MHC class I molecules.
11. The system of claim 10, wherein the peptides included in the set of training peptides have a binding affinity for mouse MHC class I K^b.
12. The system of claim 9, wherein the set of test peptides include peptides having a binding affinity for MHC class I molecules.
13. The system of claim 12, wherein the peptides included in the set of test peptides have a binding affinity for mouse MHC class I K^b.
14. The system of claims 8 or 9, wherein the ANN comprises a multi-layer perceptron ANN trained by back-propagation of error.

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15. A computer program, residing on a computer-readable medium, for identifying relative binding motifs for peptide-like molecules, comprising instructions for causing a computer to:
- (a) train an artificial neural network (ANN) with a set of training peptide-like molecules, each of known sequence and binding affinity;
 - (b) apply to the ANN at least one test peptide-like molecule, each of known sequence but unknown binding affinity;
 - (c) analyze each applied test peptide-like molecule using the ANN to predict a relative binding affinity for each test peptide-like molecule.
16. A computer program, residing on a computer-readable medium, for identifying relative peptide binding motifs, comprising instructions for causing a computer to:
- (a) train an artificial neural network (ANN) with a set of training peptides, each of known binding affinity, each peptide comprising a sequence of amino acids, each amino acid being binary coded as having or lacking specific features generally characteristic of amino acids;
 - (b) apply to the ANN at least one test peptide, each of unknown binding affinity, each peptide comprising a sequence of amino acids, each amino acid being binary coded as having or lacking specific features generally characteristic of amino acids;
 - (c) analyze each applied test peptide using the ANN to predict a relative binding affinity for each test peptide.

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17. The computer program of claim 16, wherein the set of training peptides having a binding affinity for MHC class I molecules.
18. The computer program of claim 17, wherein the peptides included in the set of training peptides have a binding affinity for mouse MHC class I K^b.
19. The computer program of claim 16, wherein the set of test peptides include peptides having a binding affinity for MHC class I molecules.
20. The computer program of claim 19, wherein the peptides included in the set of test peptides have a binding affinity for mouse MHC class I K^b.
21. The computer program of claims 15 or 16, wherein the ANN comprises a multi-layer perceptron ANN trained by back-propagation of error.

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